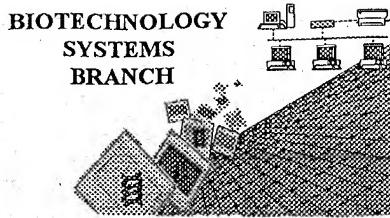


0570  
0709



## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/033,195  
Source: OIPF  
Date Processed by STIC: 7/9/2002

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
3. Hand Carry directly to:  
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7<sup>th</sup> Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202  
Or  
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002

**Raw Sequence Listing Error Summary**

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	<u>SERIAL NUMBER:</u>
		<u>10/033,195</u>

**ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE**

- 1  Wrapped Nucleic  
Wrapped Aminos
 

The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2  Invalid Line Length
 

The rules require that a line not exceed 72 characters in length. This includes white spaces.
- 3  Misaligned Amino  
Numbering
 

The numbering under each 5<sup>th</sup> amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.
- 4  Non-ASCII
 

The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
- 5  Variable Length
 

Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6  PatentIn 2.0  
"bug"
 

A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) \_\_\_\_\_. Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
- 7  Skipped Sequences  
(OLD RULES)
 

Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:  
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)  
 (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)  
 This sequence is intentionally skipped  
  
 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8  Skipped Sequences  
(NEW RULES)
 

Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.  
<210> sequence id number  
<400> sequence id number  
000
- 9  Use of n's or Xaa's  
(NEW RULES)
 

Use of n's and/or Xaa's have been detected in the Sequence Listing.  
Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.  
In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10  Invalid <213>  
Response
 

Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
- 11  Use of <220>
 

Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.  
Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." (Please explain source of genetic material in <220> to <223> section.)  
(See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12  PatentIn 2.0  
"bug"
 

Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13  Misuse of n
 

n can only be used to represent a single nucleotide in a nucleic acid sequence. N is not used to represent any value not specifically a nucleotide.

**IMPORTANT**  
*(e.g. peptide  
Sequence  
would be  
insufficient)*



OIPE

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/033,195

DATE: 07/09/2002  
TIME: 13:34:10

Input Set : A:\2719.2002-001.txt  
Output Set: N:\CRF3\07092002\J033195.raw

pp 1-4

Does Not Comply  
Corrected Diskette Needed

4 <110> APPLICANT: Fodor, Stephen P.A.  
 5 Stryer, Lubert  
 6 Read, J. Leighton  
 7 Pirrung, Michael C.  
 9 <120> TITLE OF INVENTION: Nucleotides and Analogs Having  
 10 Photoremovable Protecting Groups  
 13 <130> FILE REFERENCE: 2719.2002-001  
 15 <140> CURRENT APPLICATION NUMBER: 10/033,195  
 16 <141> CURRENT FILING DATE: 2002-06-24  
 18 <150> PRIOR APPLICATION NUMBER: 09/465,126  
 19 <151> PRIOR FILING DATE: 1999-12-17  
 21 <150> PRIOR APPLICATION NUMBER: 09/063,933  
 22 <151> PRIOR FILING DATE: 1998-04-21  
 24 <150> PRIOR APPLICATION NUMBER: 08/466,632  
 25 <151> PRIOR FILING DATE: 1995-06-06  
 27 <150> PRIOR APPLICATION NUMBER: 08/390,272  
 28 <151> PRIOR FILING DATE: 1995-02-16  
 30 <150> PRIOR APPLICATION NUMBER: 07/624,120  
 31 <151> PRIOR FILING DATE: 1990-12-06  
 33 <150> PRIOR APPLICATION NUMBER: 07/492,462  
 34 <151> PRIOR FILING DATE: 1990-03-07  
 36 <150> PRIOR APPLICATION NUMBER: 07/362,901  
 37 <151> PRIOR FILING DATE: 1989-06-07  
 39 <150> PRIOR APPLICATION NUMBER: 08/456,887  
 40 <151> PRIOR FILING DATE: 1995-06-01  
 42 <150> PRIOR APPLICATION NUMBER: 07/954,646  
 43 <151> PRIOR FILING DATE: 1992-09-30  
 45 <150> PRIOR APPLICATION NUMBER: 07/850,356  
 46 <151> PRIOR FILING DATE: 1992-03-12  
 49 <160> NUMBER OF SEQ ID NOS: 20  
 51 <170> SOFTWARE: FastSEQ for Windows Version 4.0  
 53 <210> SEQ ID NO: 1  
 54 <211> LENGTH: 5  
 55 <212> TYPE: PRT  
 56 <213> ORGANISM: Peptide sequence (global)  
 58 <400> SEQUENCE: invalid response - see item 10 on Error  
 59 Tyr Gly Gly Phe Leu  
 60 1 5  
 63 <210> SEQ ID NO: 2  
 64 <211> LENGTH: 5  
 65 <212> TYPE: PRT  
 66 <213> ORGANISM: Peptide sequence  
 68 <400> SEQUENCE: 2

summary  
sheet

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/033,195

DATE: 07/09/2002  
TIME: 13:34:10

Input Set : A:\2719.2002-001.txt  
Output Set: N:\CRF3\07092002\J033195.raw

69 Pro Gly Gly Phe Leu  
70 1 5  
73 <210> SEQ ID NO: 3  
74 <211> LENGTH: 6  
75 <212> TYPE: PRT  
76 <213> ORGANISM: Peptide sequence  
78 <400> SEQUENCE: 3  
79 Tyr Gly Ala Phe Leu Ser  
80 1 5  
83 <210> SEQ ID NO: 4  
84 <211> LENGTH: 5  
85 <212> TYPE: PRT  
86 <213> ORGANISM: Peptide sequence  
88 <400> SEQUENCE: 4  
89 Tyr Gly Ala Phe Ser  
90 1 5  
93 <210> SEQ ID NO: 5  
94 <211> LENGTH: 5  
95 <212> TYPE: PRT  
96 <213> ORGANISM: Peptide sequence  
98 <400> SEQUENCE: 5  
99 Tyr Gly Ala Phe Leu  
100 1 5  
103 <210> SEQ ID NO: 6  
104 <211> LENGTH: 6  
105 <212> TYPE: PRT  
106 <213> ORGANISM: Peptide sequence  
108 <400> SEQUENCE: 6  
109 Tyr Gly Gly Phe Leu Ser  
110 1 5  
113 <210> SEQ ID NO: 7  
114 <211> LENGTH: 4  
115 <212> TYPE: PRT  
116 <213> ORGANISM: Peptide sequence  
118 <400> SEQUENCE: 7  
119 Thr Gly Ala Phe  
120 1  
123 <210> SEQ ID NO: 8  
124 <211> LENGTH: 5  
125 <212> TYPE: PRT  
126 <213> ORGANISM: Peptide sequence  
128 <400> SEQUENCE: 8  
129 Tyr Gly Ala Leu Ser  
130 1 5  
133 <210> SEQ ID NO: 9  
134 <211> LENGTH: 5  
135 <212> TYPE: PRT  
136 <213> ORGANISM: Peptide sequence  
138 <400> SEQUENCE: 9

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/033,195

DATE: 07/09/2002  
TIME: 13:34:10

Input Set : A:\2719.2002-001.txt  
Output Set: N:\CRF3\07092002\J033195.raw

139 Tyr Gly Gly Phe Ser  
140 1 5  
143 <210> SEQ ID NO: 10  
144 <211> LENGTH: 4  
145 <212> TYPE: PRT  
146 <213> ORGANISM: Peptide sequence  
148 <400> SEQUENCE: 10  
149 Tyr Gly Ala Leu  
150 1  
153 <210> SEQ ID NO: 11  
154 <211> LENGTH: 6  
155 <212> TYPE: PRT  
156 <213> ORGANISM: Peptide sequence  
158 <400> SEQUENCE: 11  
159 Tyr Gly Ala Phe Leu Phe  
160 1 5  
163 <210> SEQ ID NO: 12  
164 <211> LENGTH: 5  
165 <212> TYPE: PRT  
166 <213> ORGANISM: Peptide sequence  
168 <400> SEQUENCE: 12  
169 Tyr Gly Ala Phe Phe  
170 1 5  
173 <210> SEQ ID NO: 13  
174 <211> LENGTH: 5  
175 <212> TYPE: PRT  
176 <213> ORGANISM: Peptide sequence  
178 <400> SEQUENCE: 13  
179 Tyr Gly Gly Leu Ser  
180 1 5  
183 <210> SEQ ID NO: 14  
184 <211> LENGTH: 6  
185 <212> TYPE: PRT  
186 <213> ORGANISM: Peptide sequence  
188 <400> SEQUENCE: 14  
189 Tyr Gly Ala Phe Ser Phe  
190 1 5  
193 <210> SEQ ID NO: 15  
194 <211> LENGTH: 7  
195 <212> TYPE: PRT  
196 <213> ORGANISM: Peptide sequence  
198 <400> SEQUENCE: 15  
199 Tyr Gly Ala Phe Leu Ser Phe  
200 1 5  
203 <210> SEQ ID NO: 16  
204 <211> LENGTH: 6  
205 <212> TYPE: PRT  
206 <213> ORGANISM: Peptide sequence  
208 <400> SEQUENCE: 16

RAW SEQUENCE LISTING  
PATENT APPLICATION: US/10/033,195

DATE: 07/09/2002  
TIME: 13:34:10

Input Set : A:\2719.2002-001.txt  
Output Set: N:\CRF3\07092002\J033195.raw

209 Tyr Gly Ala Phe Met Gln  
210 1 5  
213 <210> SEQ ID NO: 17  
214 <211> LENGTH: 5  
215 <212> TYPE: PRT  
216 <213> ORGANISM: Peptide sequence  
218 <400> SEQUENCE: 17  
219 Tyr Gly Ala Phe Met  
220 1 5  
223 <210> SEQ ID NO: 18  
224 <211> LENGTH: 5  
225 <212> TYPE: PRT  
226 <213> ORGANISM: Peptide sequence  
228 <400> SEQUENCE: 18  
229 Tyr Gly Ala Phe Gln  
230 1 5  
233 <210> SEQ ID NO: 19  
234 <211> LENGTH: 5  
235 <212> TYPE: PRT  
236 <213> ORGANISM: Peptide Sequence  
238 <400> SEQUENCE: 19  
239 Tyr Gly Gly Phe Met  
240 1 5  
243 <210> SEQ ID NO: 20  
244 <211> LENGTH: 4  
245 <212> TYPE: PRT  
246 <213> ORGANISM: Peptide sequence  
248 <400> SEQUENCE: 20  
249 Gly Gly Phe Leu  
250 1

VERIFICATION SUMMARY  
PATENT APPLICATION: US/10/033,195

DATE: 07/09/2002  
TIME: 13:34:11

Input Set : A:\2719.2002-001.txt  
Output Set: N:\CRF3\07092002\J033195.raw

L:16 M:271 C: Current Filing Date differs, Replaced Current Filing Date